TGTCAAACACACACATAACACATAAGTGACCGTGAGTCATTAAATTTATATATATTCATCAATC

TAATCAAACTATGGAGAAGAAATCACTAGCTGGCTTATGCTTCCTCTTCTTGGTTCTCTTTGTT
M E K K S L A G L C F L F L V L F V

GCACAAGAAATTATGGTGACAGAAGCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGAC
E Q E I M V T E A R T C E N L A D K Y R G P
Δ

CATGCTTTAGTGGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGTAG
C F S G C D T H C T T K E N A V S G R C R

GGACGACTTCCGCTGCTGGTGTACTAAAAGATGTTAAATGGATCTCCTCCAACATCAAGATGTG
D D F R C W C T K R C *

CAAATAGTCTTTATAATAAAACTAAATAAATAAAATGCACGCAGTATAGCTACAACTTCATCTA

TTATATGTACTCAATATCGNGCATAACGTATTAGTTATGCACTTCTATCATATGGAATAAACAT

AATAAGTAATTTCGTNTCCAAAAAAAAAAAAAAAAAAA

FIG. 1

FIG. 1

Red-lined version

```
AlfAFP2     TGTCAAACACACACATAACACATAAGTGACCGTGAGTCATTAAATTTATA
ALFAFP1     --------------------------------------------------

AlfAFP2     TATATTCATCAATCTAATCAAACTATGGAGAAGAAATCACTAGCTGGCTTA
AlfAFP1     ----------------------------------------------CAGGCTTA
                                                          * ******

AlfAFP2     TGCTTCCTCTTCCTCGTTCTCTTTGTTGAACAAGAAATTATGGTGACCGAG
AlfAFP1     TGCTTCCTCTTCTTGGTTCTCTTTGTTGCACAAGAAATTGTGGTGACAGAA
            ************ * ************** ********** ******* **

AlfAFP2     GCAGCTACTTGTGAGAATTTGGCTAACACATACAGGGGACCATGCTTCGGT
AlfAFP1     GCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGACCATGCTTTAGT
            **     ** ***************  * * *** ***************** **

AlfAFP2     GGTTGTGACTTTCACTGCAAAACCAAAGAACACTTACTTAGCGGXAGGTGC
AlfAFP1     GGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGT
            **********  ******** ********* **  * **** ** *****

AlfAFP2     AGGGACGACTTCCGCTGCTGCTGGATCC
AlfAFP1     AGGGACGACTTCCGCTGCTGCTGGATCC
            ****************************
```

FIG. 3

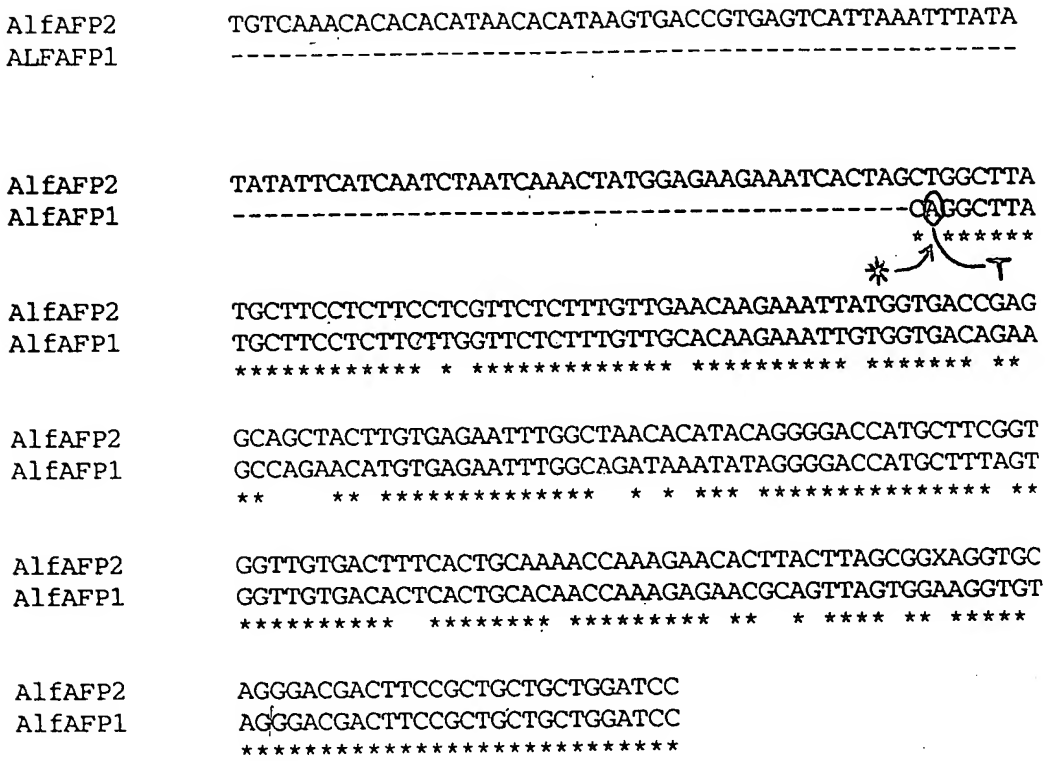

FIG. 3

Red-lined version